AMENDMENTS TO THE DRAWINGS:

The attached sheet of drawings includes a correction to Figure 1. The corrected drawing places the number 40 above the amino acid located at this position. In original Figure 1, the number 40 was placed above the amino acid located at position 39.

Attachment:

Replacement Sheet

Annotated Sheet Showing Changes

Application No. 09/976,414 Amendment Dated September 16, 2003 Replacement Sheet

NO:	1	10	20		30	40	50				
a)	AQSVPYGV	SQIKAPALHS	SQGYTO	SNVKV	AVIDSGI	DSSHPDLKVA	GGASM				
b)	AQSVPWGI:	SRVQAPAAHN	RGLTC	SGVKV	AVLDTGI	*STHPDLNIR	GGASF				
							•				
No:	`.'	60	70		80	90	100				
a)	VPSETNPFQDNNSHGTHVAGTVAALNNSIGVLGVAPSASLYAVKVLGADG										
b)	VPGEPST*QDGNGHGTHVAGTVAALNNSIGVIGVAPSADLYAVKVLGANG										
No:		110	120		130	140	150				
a)	SGQYSWIIN	GIEWA IANN	MIVOM	MSLGGF	SGSAAL	KAAVDKAVAS	GVVVV				
b)	RGSVSGIAQGLEWAAANNMHIANMSLGSDAPSTTLERAVNYATSQGVLVI										
No:		160	170		180	190	200				
a)	AAAGNEGTSGSSSTVGYPGKYPSVIAVGAVDSSNQRASFSSVGPELDVMA										
b)	AATGNNG*SGS***VGYPARYANAMAVGATDQNNRRANFSQYGTGIDIVA										
			,								
No:		210	220		230	240	250				
a)	PGVSIQSTL	PGNKYGAYN	GTSMA	3PHVAG	AAALILS	KHPNWTNTQV					
b)						KNPSWSNVQ1					
No:	:	260	270	275		•	-1-				
a)	ENTTTKLGDSFYYGKGLINVQAAAQ										
b)	KNTATSLGSTNLYGSGLVNAEAATR										

Fig. 1

Application No. 09/976,414 Amendment Dated September 16, 2003 Annotated Sheet Showing Changes

No:	: 1	10	20	30	_40	50					
a)	AQSVPYGVSQIKAPALHSQGYTGSNVKVAVIDSGIDSSHPDLKVAGGASM										
b)	AQSVPWGISRVQAPAAHNRGLTGSGVKVAVLDTGI*\$THPDLNIRGGASF										
No:		60	70	80	90	100					
a)	VPSETNPFQDNNSHGTHVAGTVAALNNSIGVLGVAPSASLYAVKVLGADG										
b _i)	VPGEPST*QDGNGHGTHVAGTVAALNNSIGVIGVAPSADLYAVKVLGANG										
No:		110	120	130	140	150					
a)	SGQYSWIINGIEWAIANNMDVINMSLGGPSGSAALKAAVDKAVASGVVVV										
þ)	RGSVSGIAQGLEWAAANNMHIANMSLGSDAPSTTLERAVNYATSQGVLVI										
No:		160	170	180	190	200					
a)	AAAGNEGTSGSSSTVGYPGKYPSVIAVGAVDSSNQRASFSSVGPELDVMA										
b)	AATGNNG*SGS***VGYPARYANAMAVGATDQNNRRANFSQYGTGIDIVA										
No:	7	210	220	230	240	250					
a)	PGVSIQSTLPGNKYGAYNGTSMASPHVAGAAALILSKHPNWTNTQVRSSL										
b)	PGVDIESTYPGSSYDSLSGTSMATPHVAGVAALVKQKNPSWSNVQIRNHL										
No:	2	60	270 275								
a)	ENTTTKLGDSFYYGKGLINVQAAAQ										
þ)	KNTATSLGSTNLYGSGLVNAEAATR										

Fig. 1